STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/557,992_
Source: _PCT/10_
Date Processed by STIC: _12/2/05_

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PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
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Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**
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Revised 01/24/05

PCT

RAW SEQUENCE LISTING   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:25

Input Set : **A:\PTO.SR.txt**
Output Set: **N:\CRF4\12022005\J557992.raw**

```
     3 <110> APPLICANT: Japan Science and Technology Corporation
     5 <120> TITLE OF INVENTION: Probe for visualizing protein interaction and method of
analyzing protein-
     6        protein interaction using the same
     8 <130> FILE REFERENCE: 04F025PCT
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/557,992
C--> 10 <141> CURRENT FILING DATE: 2005-11-22
E--> 10 <160> NUMBER OF SEQ ID NOS: 7 (p.4)
```

Handwritten: insert <140> 10/557,992 <141> 2005-11-22

Handwritten: Does Not Comply Corrected Diskette Needed see pp 1-6

ERRORED SEQUENCES

```
    12 <210> SEQ ID NO: 1
    13 <211> LENGTH: 300
    14 <212> TYPE: PRT
    15 <213> ORGANISM: Renilla reniformis
W--> 16 <400> SEQUENCE: 1
    17     Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile
    18     1               5                   10                  15
    19     Thr Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu
E--> 20     16              20                  25                  30
    21     Asp Ser Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn
E--> 22     31              35                  40                  45
    23     Ala Val Ile Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp
E--> 24     46              50                  55                  60
    25     Arg His Val Val Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile
E--> 26     61              65                  70                  75
    27     Pro Asp Leu Ile Gly Met Gly Lys Ser Gly Lys Ser Gly Asn Gly
E--> 28     76              80                  85                  90
    29     Ser Tyr Arg Leu Leu Asp His Tyr Lys Tyr Leu Thr Ala Trp Phe
E--> 30     91              95                  100                 105
    31     Glu Leu Leu Asn Leu Pro Lys Lys Ile Ile Phe Val Gly His Asp
E--> 32     106             110                 115                 120
    33     Trp Gly Ala Cys Leu Ala Phe His Tyr Cys Tyr Glu His Gln Asp
E--> 34     121             125                 130                 135
    35     Lys Ile Lys Ala Ile Val His Ala Glu Ser Val Val Asp Val Ile
E--> 36     136             140                 145                 150
    37     Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu Glu Asp Ile Ala Leu
E--> 38     151             155                 160                 165
    39     Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu Glu Asn Asn Phe
E--> 40     166             170                 175                 180
    41     Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg Lys Leu Glu
E--> 42     181             185                 190                 195
    43     Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu Lys Gly
```

Handwritten margin notes: Delete these nos; Per 1.822 of Sequence Rules, number the amino acids under every — 5 amino = acids only; see p. 2

```
                    RAW SEQUENCE LISTING              DATE: 12/02/2005
                    PATENT APPLICATION:  US/10/557,992    TIME: 11:33:25

                    Input Set : A:\PTO.SR.txt
                    Output Set: N:\CRF4\12022005\J557992.raw
```

```
E--> 44    196                200                205                210
     45  Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro Leu
E--> 46    211                215                220                225
     47  Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn Tyr
E--> 48    226                230                235                240
     49  Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe Ile
E--> 50    241                245                250                255
     51  Glu Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly Ala
E--> 52    256                260                265                270
     53  Lys Lys Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu His
E--> 54    271                275                280                285
     55  Phe Ser Gln Glu Asp Ala Pro Asp Glu Met Gly Asn Tyr Ile Gln
E--> 56    286            .    290→             295→               300→
     59  <210> SEQ ID NO: 2
     60  <211> LENGTH: 13
     61  <212> TYPE: PRT
     62  <213> ORGANISM: Artificial Sequence
W--> 63  <220>
     63  <223> OTHER INFORMATION: Synthesized Oligopeptide
     64  <400> SEQUENCE: 2
     65    Thr Glu Glu Ala Tyr Met Lys Met Asp Leu Gly Pro Gly
     66    1                5                10
     69  <210> SEQ ID NO: 3
     70  <211> LENGTH: 300
     71  <212> TYPE: PRT
     72  <213> ORGANISM: Renilla reniformis
W--> 73  <400> SEQUENCE: 3
     74    Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile
     75    1                5                10                15
     76    Thr Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu
E--> 77    16               20                25                30
     78    Asp Ser Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn
E--> 79    31               35                40                45
     80    Ala Val Ile Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp
E--> 81    46               50                55                60
     82    Arg His Val Val Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile
E--> 83    61               65                70                75
     84    Pro Asp Leu Ile Gly Met Gly Lys Ser Gly Lys Ser Gly Asn Gly
E--> 85    76               80                85                90
     86    Ser Tyr Arg Leu Leu Asp His Tyr Lys Tyr Leu Thr Ala Trp Phe
E--> 87    91               95                100               105
     88    Glu Leu Leu Asn Leu Pro Lys Lys Ile Ile Phe Val Gly His Asp
E--> 89    106              110               115               120
     90    Trp Gly Ala Ala Leu Ala Phe His Tyr Cys Tyr Glu His Gln Asp
E--> 91    121              125               130               135
     92    Lys Ile Lys Ala Ile Val His Ala Glu Ser Val Val Asp Val Ile
E--> 93    136              140               145               150
     94    Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu Glu Asp Ile Ala Leu
E--> 95    151              155               160               165
```

[Handwritten annotations:]

290→ directly under "Asp" 295→ directly under "Met" 300→ directly under "Gln"

<220> ← insert <220> whenever <221>, <222>, or <223> is shown. <220> never has a response. It is a "header" only.

same error as sequence 1

RAW SEQUENCE LISTING   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:25

Input Set : A:\PTO.SR.txt
Output Set: N:\CRF4\12022005\J557992.raw

```
    96    Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu Glu Asn Asn Phe
E--> 97    166         170             175             180
    98    Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg Lys Leu Glu
E--> 99    181         185             190             195
   100     Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu Lys Gly
E--> 101   196         200             205             210
   102     Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro Leu
E--> 103   211         215             220             225
   104     Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn Tyr
E--> 105   226         230             235             240
   106     Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe Ile
E--> 107   241         245             250             255
   108     Glu Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly Ala
E--> 109   256         260             265             270
   110     Lys Lys Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu His
E--> 111   271         275             280             285
   112     Phe Ser Gln Glu Asp Ala Pro Asp Glu Met Gly Asn Tyr Ile Gln
   113     286         290             295             300
   116 <210> SEQ ID NO: 4
   117 <211> LENGTH: 18
   118 <212> TYPE: PRT
   119 <213> ORGANISM: Artificial Sequence
W--> 120 <220>
   120 <223> OTHER INFORMATION: Synthesized Oligopeptide
   121 <400> SEQUENCE: 4
   122     Cys Leu Ser Leu Ala Ser Asn Asn Gly Asn Gly Arg Asn Gly Ala
   123     1               5               10              15
   124     Ser Leu Glu
   125     16
   128 <210> SEQ ID NO: 5
   129 <211> LENGTH: 17
   130 <212> TYPE: PRT
   131 <213> ORGANISM: Artificial Sequence
W--> 132 <220>
   132 <223> OTHER INFORMATION: Synthesized Oligopeptide
   133 <400> SEQUENCE: 5
   134     Pro Arg Gly Asn Asn Gly Gly Asn Asn Asp Val Met Ala Ile Ala
   135     1               5               10              15
   136     Ala Asn
   137     16
   140 <210> SEQ ID NO: 6
   141 <211> LENGTH: 133
   142 <212> TYPE: PRT
   143 <213> ORGANISM: Artificial Sequence
W--> 144 <220>
   144 <223> OTHER INFORMATION: Synthesized Oligopeptide
   145 <400> SEQUENCE: 6
   146     Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile
   147     1               5               10              15
```

handwritten annotations: directly "under Met" (at line 113, near 295)

handwritten annotations: ← insert <220> (at lines 120, 132, 144)

handwritten annotations: delete (at lines 125, 137)

RAW SEQUENCE LISTING   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:25

Input Set : A:\PTO.SR.txt
Output Set: N:\CRF4\12022005\J557992.raw

```
        148     Thr Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu
E--> 149        16          20              25              30
        150     Asp Ser Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn
E--> 151        31          35              40              45
        152     Ala Val Ile Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp
E--> 153        46          50              55              60
        154     Arg His Val Val Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile
E--> 155        61          65              70              75
        156     Pro Asp Leu Ile Gly Met Gly Lys Ser Gly Lys Ser Gly Asn Gly
E--> 157        76          80              85              90
        158     Ser Cys Leu Ser Leu Ala Ser Asn Asn Gly Asn Gly Arg Asn Gly
E--> 159        91          95              100             105
        160     Ala Ser Leu Glu Thr Glu Glu Tyr Met Lys Met Asp Leu Gly Pro
E--> 161        106         110             115             120
        162     Gly Thr Arg Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
E--> 163        121         125->           130->
        166 <210> SEQ ID NO: 7
        167 <211> LENGTH: 352
        168 <212> TYPE: PRT
        169 <213> ORGANISM: Artificial Sequence
W--> 170 <220>
        170 <223> OTHER INFORMATION: Synthesized Oligopeptide
        171 <400> SEQUENCE: 7
        172     Met Asp Ala Glu Trp Tyr Trp Gly Asp Ile Ser Arg Glu Glu Val
        173     1           5               10              15
        174     Asn Glu Lys Leu Arg Asp Thr Ala Asp Gly Thr Phe Leu Val Arg
E--> 175        16          20              25              30
        176     Asp Ala Ser Thr Lys Met His Gly Asp Tyr Thr Leu Thr Leu Arg
E--> 177        31          35              40              45
        178     Lys Gly Gly Asn Asn Lys Leu Ile Lys Ile Phe His Arg Asp Gly
E--> 179        46          50              55              60
        180     Lys Tyr Gly Phe Ser Asp Pro Leu Thr Phe Asn Ser Val Val Glu
E--> 181        61          65              70              75
        182     Leu Ile Asn His Tyr Arg Asn Glu Ser Leu Ala Gln Tyr Asn Pro
E--> 183        76          80              85              90
        184     Lys Leu Asp Val Lys Leu Leu Tyr Pro Val Ser Lys Tyr Gln Gln
E--> 185        91          95              100             105
        186     Pro Arg Gly Asn Asn Gly Gly Asn Asn Asp Val Met Ala Ile Ala
E--> 187        106         110             115             120
        188     Ala Asn Tyr Arg Leu Leu Asp His Tyr Lys Tyr Leu Thr Ala Trp
E--> 189        121         125             130             135
        190     Phe Glu Leu Leu Asn Leu Pro Lys Lys Ile Ile Phe Val Gly His
E--> 191        136         140             145             150
        192     Asp Trp Gly Ala Cys Leu Ala Phe His Tyr Ser Tyr Glu His Gln
E--> 193        151         155             160             165
        194     Asp Lys Ile Lys Ala Ile Val His Ala Glu Ser Val Val Asp Val
E--> 195        166         170             175             180
        196     Ile Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu Glu Asp Ile Ala
E--> 197        181         185             190             195
```

[Handwritten annotations:] 125→ directly under Gln; 130→ directly under "Glu"; last sequence in submitted file;

RAW SEQUENCE LISTING   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:25

Input Set : A:\PTO.SR.txt
Output Set: N:\CRF4\12022005\J557992.raw

```
        198    Leu Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu Glu Asn Asn
E--> 199    196             200             ·             205                     210
        200    Phe Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg Lys Leu
E--> 201    211             215                         220                     225
        202    Glu Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu Lys
E--> 203    226             230                         235                     240
        204    Gly Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro
E--> 205    241             245                         250                     255
        206    Leu Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn
E--> 207    256             260                         265                     270
        208    Tyr Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe
E--> 209    271             275                         280                     285
        210    Ile Glu Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly
E--> 211    286             290                         295                     300
        212    Ala Lys Lys Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu
E--> 213    301             305                         310                     315
        214    His Phe Ser Gln Glu Asp Ala Pro Asp Glu Met Gly Lys Tyr Ile
E--> 215    316             320                         325                     330
        216    Lys Ser Phe Val Glu Arg Val Leu Lys Asn Glu Gln Pro Arg Asp
E--> 217    331             335                         340                     345
        218    Tyr Lys Asp Asp Val Val Lys
        219    346             350
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 12/02/2005
PATENT APPLICATION: **US/10/557,992**   TIME: 11:33:26

Input Set : **A:\PTO.SR.txt**
Output Set: **N:\CRF4\12022005\J557992.raw**

<u>**Invalid Line Length:**</u>
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

VERIFICATION SUMMARY   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:26

Input Set : **A:\PTO.SR.txt**
Output Set: **N:\CRF4\12022005\J557992.raw**

```
L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:16 M:283 W: Missing Blank Line separator, <400> field identifier
L:20 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:22 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:24 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:26 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:28 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:30 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:32 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:34 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:36 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:38 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:40 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:42 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:44 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:46 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:48 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:50 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:52 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:54 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:56 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:63 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
L:64 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:2
L:64 M:283 W: Missing Blank Line separator, <400> field identifier
L:73 M:283 W: Missing Blank Line separator, <400> field identifier
L:77 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:79 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:81 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:83 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:85 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:87 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:89 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:91 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:93 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:95 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:97 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:99 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:101 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:103 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:105 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:107 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:109 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:111 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:120 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:121 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:4
L:121 M:283 W: Missing Blank Line separator, <400> field identifier
L:132 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
```

VERIFICATION SUMMARY   DATE: 12/02/2005
PATENT APPLICATION: **US/10/557,992**   TIME: 11:33:26

Input Set : **A:\PTO.SR.txt**
Output Set: **N:\CRF4\12022005\J557992.raw**

```
L:133 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:5
L:133 M:283 W: Missing Blank Line separator, <400> field identifier
L:144 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:145 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:6
L:145 M:283 W: Missing Blank Line separator, <400> field identifier
L:149 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:151 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:153 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:155 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:157 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:159 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:161 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:163 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:170 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:171 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:7
L:171 M:283 W: Missing Blank Line separator, <400> field identifier
L:175 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:177 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:179 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:181 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:183 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:185 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:187 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:189 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:191 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:193 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:195 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:197 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:199 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:201 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:203 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:205 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:207 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:209 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:211 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:213 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:215 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:217 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:10 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (6) Counted (7)
```